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Supplemental Information

Programmed Secretion Arrest and Receptor-Triggered Toxin Export during Antibacterial Contact-Dependent Growth Inhibition

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Table S1. Tomograms collected, Related to Figures 1 and 3.

Strain	Plasmid	Sample	Tomograms with filaments	Total tomograms collected
EC93 *bamA*^{LT2}	none	bacteria	21	21
EC93 Δ*cdiA*	none	bacteria	0	16
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pDAL660Δ1-39 CdiA^{EC93}	bacteria	21	21
EPI100 Δ*bamA::cat* pZS21-*bamA*$^{\mathbf{ECL}}$	pDAL660Δ1-39 CdiA^{EC93}	minicells	24	24
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pDAL660Δ1-39 CdiA$^{\mathbf{EC93}}$	bacteria + solubilized BamA	3	3
EPI100 Δ*bamA::cat* pZS21-b*amA*^{ECL}	pDAL660Δ1-39 CdiA$^{\mathbf{EC93}}$	minicells + buffer/detergent	6	6
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pDAL660Δ1-39 CdiA$^{\mathbf{EC93}}$	minicells + solubilized BamA	38	38
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pDAL660Δ1-39 CdiA$^{\mathbf{EC93}}$	minicells + target minicells (vesicles)	42	42
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pWEB-TNC (no CdiA)	minicells	0	11
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pCH12707 CdiA(Gly1318)^{EC93}	minicells	0	17
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pCH12705 CdiA(Thr2122)^{EC93}	minicells	14	14
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pCH12706 CdiA(Val1929)^{EC93}	minicells	10	10
EPI100 Δ*bamA::cat* pZS21-*bamA*^{ECL}	pZR113 CdiA(Ser832)^{EC93}	minicells	0	13
MG1655 Δ*wzb* Δ*tsx* Δ*ompT*	pET21b (no CdiA)	bacteria	0	11
MG1655 Δ*wzb* Δ*tsx* Δ*ompT*	pET21b (no CdiA)	minicells	0	1
MG1655 Δ*wzb* Δ*tsx* Δ*ompT*	pCH13604 CdiA^{STECO31}	bacteria	14	14
MG1655 Δ*wzb* Δ*tsx* Δ*ompT*	pCH13604 CdiA^{STECO31}	minicells	23	23

Table S2. CdiA domain analysis, Related to Figures 7B and 7C.

			FHA-1 region			RBD[a]			YP domain			FHA-2 domain[b]		
NCBI identifier	**Bacterial species and strain**	**CdiA length**	**begin**	**end**	**length**	**begin**	**end**	**length**	**begin**	**end**	**length**	**begin**	**end**	**length**
WP_109054141.1	*Brenneria roseae* subsp. americana	**5237**	392	3171	**2780**	3172	3619	**448**	3620	3805	**186**	3979	4656	**678**
WP_095834043.1	***Brenneria goodwinii* FRB141**	**5550**	477	3114	**2638**	3115	3562	**448**	3563	3748	**186**	4054	4863	**810**
WP_053085546.1	*Brenneria goodwinii* OBR1	**5538**	477	3114	**2638**	3115	3552	**438**	3553	3738	**186**	4044	4853	**810**
WP_064541030.1	*Buttiauxella ferragutiae* ATCC 51602	**3961**	312	2025	**1714**	2026	2362	**337**	2363	2547	**185**	2717	3321	**605**
WP_051880442.1	*Buttiauxella agrestis* MCE	**3828**	311	1896	**1586**	1897	2220	**324**	2221	2405	**185**	2420	3179	**760**
WP_039294138.1	*Cedecea neteri* M006	**5018**	310	2694	**2385**	2695	3094	**400**	3095	3279	**185**	3448	4320	**873**
WP_038482520.1	*Cedecea neteri* SSMD04	**4373**	310	2383	**2074**	2384	2790	**407**	2791	2975	**185**	3144	3689	**546**
WP_061277518.1	***Cedecea neteri* NBRC 105707**	**4616**	310	2696	**2387**	2697	3114	**418**	3115	3299	**185**	3468	4013	**546**
WP_039303703.1	*Cedecea neteri* ND14a	**4668**	310	2696	**2387**	2697	3111	**415**	3112	3296	**185**	3326	4073	**748**
WP_016151891.1	*Citrobacter* sp. KTE151	**3318**	309	1386	**1078**	1387	1662	**276**	1663	1847	**185**	2072	2625	**554**
WP_052406297.1	*Citrobacter sedlakii* NBRC 105722	**3142**	309	1262	**954**	1263	1557	**295**	1558	1742	**185**	1897	2519	**623**
WP_052443881.1	*Citrobacter* sp. JT3	**3602**	310	1357	**1048**	1358	1660	**303**	1661	1845	**185**	2005	2948	**944**
WP_102602775.1	*Citrobacter freundii* complex sp. CFNIH2	**3629**	310	1357	**1048**	1358	1660	**303**	1661	1845	**185**	2005	2948	**944**
WP_049042230.1	*Citrobacter braakii* 989_CKOS	**3950**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2639	3305	**667**
WP_016155806.1	Citrobacter sp. KTE32	**3950**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2707	3379	**673**
WP_052746947.1	*Citrobacter amalonaticus* Y19	**3628**	310	1357	**1048**	1358	1657	**300**	1658	1842	**185**	1997	2952	**956**
WP_109017325.1	*Citrobacter freundii* MH16-522D	**3950**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2707	3379	**673**
WP_105242179.1	Citrobacter sp. FDAARGOS_156	**3985**	352	2002	**1651**	2003	2346	**344**	2347	2531	**185**	2693	3308	**616**
WP_080625654.1	*Citrobacter braakii* FDAARGOS_253	**3950**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2697	3379	**683**
WP_047413132.1	*Citrobacter freundii* FDAARGOS_61	**3950**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2697	3379	**683**
WP_087051660.1	*Citrobacter europaeus* A121	**3950**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2697	3379	**683**
WP_043015339.1	*Citrobacter freundii* MGH 56	**3880**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2707	3379	**673**
WP_087053097.1	*Citrobacter europaeus* 97/99	**3950**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2697	3379	**683**
WP_085047670.1	*Citrobacter werkmanii* AK-8	**3929**	309	2001	**1693**	2002	2345	**344**	2346	2530	**185**	2707	3307	**601**
WP_103283520.1	*Citrobacter freundii* complex sp. CFNIH3	**3962**	330	2001	**1672**	2002	2357	**356**	2358	2542	**185**	2719	3391	**673**
WP_103765923.1	*Citrobacter braakii* S1283	**3825**	309	2001	**1693**	2002	2329	**328**	2330	2514	**185**	2691	3287	**597**
WP_101700591.1	*Citrobacter* sp. L55	**3901**	309	2001	**1693**	2002	2329	**328**	2330	2514	**185**	2691	3363	**673**
WP_075846765.1	*Citrobacter braakii* SCC4	**3934**	309	2001	**1693**	2002	2329	**328**	2330	2514	**185**	2691	3363	**673**
WP_016151957.1	*Citrobacter* sp. KTE151	**3901**	309	2001	**1693**	2002	2329	**328**	2330	2514	**185**	2691	3289	**599**
WP_019076112.1	*Citrobacter freundii* str. Ballerup 7851	**3901**	309	2001	**1693**	2002	2329	**328**	2330	2514	**185**	2691	3287	**597**
WP_100194023.1	*Citrobacter freundii* CRCB-101	**3962**	309	2001	**1693**	2002	2357	**356**	2358	2542	**185**	2709	3391	**683**
WP_023639573.1	*Dickeya zeae* NCPPB 3532	**3008**	307	1104	**798**	1105	1409	**305**	1410	1594	**185**	1705	2304	**600**
WP_019845538.1	*Dickeya zeae* MS1	**2975**	307	1104	**798**	1105	1392	**288**	1393	1577	**185**	1743	2355	**613**
WP_046830404.1	*Dickeya dianthicola* RNS04.9	**3839**	288	1906	**1619**	1907	2278	**372**	2279	2463	**185**	2626	3240	**615**
WP_050568668.1	*Dickeya zeae* NCPPB 2538	**3222**	307	1104	**798**	1105	1421	**317**	1422	1607	**186**	1844	2652	**809**
AAN38708.1	***Dickeya chrysanthemi* EC16**	**3848**	512	1901	**1390**	1902	2263	**362**	2264	2449	**186**	2551	3172	**622**
WP_081251380.1	*Dickeya solani* IPO 2222	**4165**	309	1933	**1625**	1934	2305	**372**	2306	2491	**186**	2599	3536	**938**
WP_013318031.1	*Dickeya dadantii* 3937	**4148**	320	2116	**1797**	2117	2480	**364**	2481	2666	**186**	2776	3391	**616**
ACS85791.1	*Dickeya paradisiaca* Ech703	**3862**	512	1904	**1393**	1905	2270	**366**	2271	2454	**184**	2617	3230	**614**
WP_107759469.1	*Dickeya* sp. Secpp 1600	**4339**	437	2061	**1625**	2062	2428	**367**	2429	2614	**186**	2724	3661	**938**
WP_100849600.1	*Dickeya fangzhongdai* DSM 101947	**4031**	437	2059	**1623**	2060	2427	**368**	2428	2613	**186**	2721	3336	**616**
WP_071842415.1	*Dickeya solani* ND14b	**4231**	438	2059	**1622**	2060	2432	**373**	2433	2618	**186**	2726	3663	**938**
WP_100849620.1	*Dickeya fangzhongdai* DSM 101947	**4056**	437	2042	**1606**	2043	2410	**368**	2411	2596	**186**	2704	3319	**616**
WP_101744646.1	*Enterobacter* sp. Crenshaw	**4465**	311	2162	**1852**	2163	2487	**325**	2488	2673	**186**	2832	3779	**948**
WP_084832595.1	*Enterobacter cloacae* UCI 39	**4142**	311	2163	**1853**	2164	2488	**325**	2489	2674	**186**	2833	3449	**617**
WP_011915321.1	*Enterobacter* sp. 638	**3967**	311	2120	**1810**	2121	2445	**325**	2446	2631	**186**	2970	3407	**438**
WP_106994740.1	*Enterobacter* sp. FS01	**3002**	309	1212	**904**	1213	1504	**292**	1505	1689	**185**	1850	2399	**550**
KDF47155.1	*Enterobacter cloacae* BIDMC 67	**3195**	310	1495	**1186**	1496	1768	**273**	1769	1953	**185**	2056	2660	**605**
WP_062856360.1	*Enterobacter* sp. RC4	**3292**	309	1495	**1187**	1496	1768	**273**	1769	1953	**185**	2047	2663	**617**
ESN17358.1	*Enterobacter* sp. MGH 24	**3243**	309	1535	**1227**	1536	1808	**273**	1809	1993	**185**	2096	2700	**605**
WP_006812043.1	*Enterobacter hormaechei* ATCC 49162	**3436**	684	1536	**853**	1537	1829	**293**	1830	2014	**185**	2247	2791	**545**
WP_084832630.1	***Enterobacter cloacae* MGH 54**	**4214**	311	2163	**1853**	2164	2505	**342**	2506	2690	**185**	2850	3466	**617**
WP_084833443.1	*Enterobacter cloacae* e2161	**4154**	311	2163	**1853**	2164	2500	**337**	2501	2686	**186**	2845	3461	**617**

WP_096179245.1	*Enterobacter asburiae* TUM10688	**4043**	273	2162	**1890**	2163	2498	**336**	2499	2684	**186**	2843	3459	**617**
WP_046091940.1	*Enterobacter cloacae* MNCRE16	**4225**	311	2163	**1853**	2164	2500	**337**	2501	2686	**186**	2845	3466	**622**
WP_073980716.1	*Enterobacter ludwigii* EnVs6	**3500**	274	1708	**1435**	1709	2015	**307**	2016	2199	**184**	2422	2907	**486**
WP_084832228.1	*Enterobacter cloacae* complex sp. 20432	**4209**	311	2163	**1853**	2164	2500	**337**	2501	2686	**186**	2845	3461	**617**
WP_084833300.1	*Enterobacter cloacae* DSM 16690	**4381**	311	2163	**1853**	2164	2499	**336**	2500	2685	**186**	2844	3791	**948**
WP_023345176.1	*Enterobacter* sp. MGH 16	**4214**	311	2163	**1853**	2164	2505	**342**	2506	2691	**186**	2850	3466	**617**
WP_001385946.1	***Escherichia coli* STEC_O31**	**3253**	310	1384	**1075**	1385	1656	**272**	1657	1841	**185**	1996	2548	**553**
AAZ57198.1	***Escherichia coli* EC93**	**3132**	310	1378	**1069**	1379	1635	**257**	1636	1820	**185**	2050	2526	**477**
WP_029589770.1	*Franconibacter helveticus* LMB 23732	**4374**	311	2363	**2053**	2364	2709	**346**	2710	2915	**206**	3137	3695	**559**
WP_073859864.1	*Franconibacter pulveris* DJ34	**4171**	312	2363	**2052**	2364	2689	**326**	2690	2895	**206**	3061	3678	**618**
WP_106930237.1	*Kluyvera ascorbata* KA2	**3302**	309	1386	**1078**	1387	1670	**284**	1671	1855	**185**	2017	2707	**691**
WP_064567413.1	*Kosakonia oryzae* Ola 51	**3984**	310	1933	**1624**	1934	2265	**332**	2266	2450	**185**	2482	3235	**754**
WP_074775709.1	*Kosakonia sacchari* SP1	**3980**	311	1644	**1334**	1645	1978	**334**	1979	2163	**185**	2195	3272	**1078**
SKC22705.1	*Kosakonia oryzae* D4	**3665**	310	1759	**1450**	1760	2091	**332**	2092	2276	**185**	2369	3061	**693**
WP_107146251.1	*Kosakonia* sp. H7A	**3671**	310	1759	**1450**	1760	2097	**338**	2098	2282	**185**	2314	3067	**754**
KDE37341.1	*Kosakonia radicincitans* UMEnt01/12	**3671**	310	1759	**1450**	1760	2097	**338**	2098	2282	**185**	2314	3067	**754**
WP_083565982.1	*Kosakonia radicincitans* GXGL-4A	**3705**	310	1759	**1450**	1760	2131	**372**	2132	2316	**185**	2348	3101	**754**
WP_052502032.1	*Kosakonia radicincitans* YD4	**4124**	310	1759	**1450**	1760	2131	**372**	2132	2316	**185**	2427	3425	**999**
WP_094419452.1	*Kosakonia cowanii* Esp_Z	**4959**	310	2850	**2541**	2851	3186	**336**	3187	3458	**272**	3628	4242	**615**
WP_082142181.1	*Leclercia* sp. LK8	**4141**	311	2163	**1853**	2164	2488	**325**	2489	2674	**186**	2833	3449	**617**
WP_082019608.1	*Leclercia adecarboxylata* KY2	**3024**	312	1251	**940**	1252	1524	**273**	1521	1706	**186**	1862	2411	**550**
WP_036108049.1	*Mangrovibacter* sp. MFB070	**3577**	311	1709	**1399**	1710	2033	**324**	2034	2219	**186**	2303	3003	**701**
WP_083581958.1	*Pantoea sesami* Si-M154	**3304**	310	1496	**1187**	1497	1769	**273**	1770	1954	**185**	2057	2661	**605**
WP_052696968.1	*Pantoea* sp. SM3	**4171**	311	1941	**1631**	1942	2278	**337**	2279	2462	**184**	2615	3483	**869**
WP_109718305.1	*Pantoea allii* PNA 200-10	**4266**	312	2033	**1722**	2034	2361	**328**	2362	2545	**184**	2707	3629	**923**
WP_105077596.1	*Pantoea ananatis* PANS 99-23	**3976**	311	2033	**1723**	2034	2361	**328**	2362	2545	**184**	2708	3320	**613**
WP_081138203.1	*Pantoea latae* AS1	**3884**	394	1935	**1542**	1936	2261	**326**	2262	2445	**184**	2614	3225	**612**
WP_081049483.1	*Pantoea stewartii* RSA30	**4032**	311	2033	**1723**	2034	2361	**328**	2362	2545	**184**	2546	3320	**775**
WP_014593028.1	*Pantoea ananatis* AJ13355	**3728**	314	1764	**1451**	1765	2113	**349**	2114	2297	**184**	2298	3072	**775**
WP_013024799.1	*Pantoea ananatis* LMG 20103	**3667**	311	1761	**1451**	1762	2091	**330**	2092	2275	**184**	2276	3055	**780**
SFN23123.1	***Pantoea* sp. OV426**	**4089**	312	1826	**1515**	1827	2161	**335**	2162	2345	**184**	2442	3445	**1004**
WP_080752852.1	*Pantoea stewartii* M009	**4046**	311	2033	**1723**	2034	2326	**293**	2327	2510	**184**	2603	3290	**688**
WP_081316794.1	*Pantoea stewartii* NS381	**4322**	311	2033	**1723**	2034	2382	**349**	2383	2566	**184**	2567	3653	**1087**
WP_085067788.1	*Pantoea alhagi* LTYR-11Z	**3779**	312	1728	**1417**	1729	2065	**337**	2066	2251	**186**	2422	3030	**609**
WP_081325946.1	*Pantoea agglomerans* C410P1	**4169**	311	1865	**1555**	1866	2212	**347**	2213	2398	**186**	2781	3496	**716**
WP_082787945.1	*Pantoea vagans* FaVv11	**4150**	311	1865	**1555**	1866	2193	**328**	2194	2378	**185**	2562	3477	**916**
WP_013358311.1	*Pantoea vagans* C9-1	**4320**	311	1978	**1668**	1979	2319	**341**	2320	2504	**185**	2679	3603	**925**
WP_104093316.1	*Pantoea* sp. ICBG 985	**3571**	275	1614	**1340**	1615	1914	**300**	1915	2116	**202**	2231	2893	**663**
WP_088517282.1	*Pantoea* sp. VS1	**3515**	275	1614	**1340**	1615	1905	**291**	1906	2107	**202**	2279	2884	**606**
POY59994.1	***Pectobacterium maceratum* PB70**	**5651**	464	3561	**3098**	3562	4052	**491**	4053	4238	**186**	4405	5014	**610**
WP_072034242.1	*Pectobacterium carotovorum* M022	**5051**	495	2967	**2473**	2968	3406	**439**	3407	3592	**186**	3759	4368	**610**
WP_014915528.1	*Pectobacterium carotovorum* PCC21	**5819**	292	3696	**3405**	3697	4196	**500**	4197	4382	**186**	4556	5158	**603**
WP_014915291.1	*Pectobacterium carotovorum* PCC21	**6075**	292	3660	**3369**	3661	4149	**489**	4150	4335	**186**	4509	5445	**937**
WP_095994655.1	*Pectobacterium polaris* NIBIO1006	**6217**	292	3749	**3458**	3750	4244	**495**	4245	4430	**186**	4597	5540	**944**
SHG12251.1	*Pectobacterium carotovorum* DSM 30168	**5857**	496	3668	**3173**	3669	4164	**496**	4165	4350	**186**	4517	5126	**610**
PPE62299.1	*Pectobacterium carotovorum* F157	**5839**	288	3666	**3379**	3667	4142	**476**	4143	4328	**186**	4502	5104	**603**
PPE64673.1	***Pectobacterium carotovorum* F152**	**6442**	496	4303	**3808**	4304	4853	**550**	4854	5039	**186**	5213	5815	**603**
AVT58658.1	*Pectobacterium carotovorum* 3-2	**5749**	519	3673	**3155**	3674	4168	**495**	4169	4354	**186**	4528	5130	**603**
POY56417.1	*Pectobacterium maceratum* F018	**5747**	520	3654	**3135**	3655	4148	**494**	4149	4334	**186**	4508	5110	**603**
WP_080735406.1	*Pectobacterium carotovorum* NCPPB 3839	**5845**	500	3669	**3170**	3670	4181	**512**	4182	4367	**186**	4534	5143	**610**
PVY74713.1	*Pectobacterium carotovorum* WS2152	**5889**	500	3710	**3211**	3711	4196	**486**	4197	4382	**186**	4550	5158	**609**
WP_094369960.1	*Pectobacterium carotovorum* BF45	**5785**	292	3667	**3376**	3668	4162	**495**	4163	4348	**186**	4522	5124	**603**
WP_080768449.1	*Pectobacterium carotovorum* YC D64	**6059**	287	3650	**3364**	3651	4136	**486**	4137	4322	**186**	4496	5432	**937**
AIK14231.1	*Pectobacterium atrosepticum* 21A	**5753**	472	3681	**3210**	3682	4157	**476**	4158	4343	**186**	4458	5119	**662**
WP_025920092.1	*Pectobacterium parmentieri* CFIA1002	**5448**	289	3030	**2742**	3031	3474	**444**	3475	3660	**186**	3834	4766	**933**

ACX88282.1	***Pectobacterium parmentieri* WPP163**	**5981**	289	3630	**3342**	3631	4106	**476**	4107	4292	**186**	4407	5397	**991**
WP_074824325.1	*Pragia fontium* ATCC 49100	**3121**	335	947	**613**	948	1195	**248**	1196	1380	**185**	1549	2435	**887**
WP_013575347.1	*Rahnella* sp. Y9602	**3756**	312	1813	**1502**	1814	2101	**288**	2102	2327	**226**	2485	3102	**618**
WP_015689887.1	*Rahnella aquatilis* HX2	**3756**	312	1813	**1502**	1814	2101	**288**	2102	2327	**226**	2485	3102	**618**
AEX52032.1	*Rahnella aquatilis* ATCC 33071	**3895**	312	1953	**1642**	1954	2243	**290**	2244	2469	**226**	2627	3246	**620**
WP_086935272.1	*Rahnella aquatilis* ATCC 33071	**3901**	318	1959	**1642**	1960	2249	**290**	2250	2475	**226**	2633	3252	**620**
WP_101079777.1	*Rahnella* sp. AA	**3431**	279	1594	**1316**	1595	1884	**290**	1885	2082	**198**	2240	2860	**621**
WP_082001717.1	*Rahnella aquatilis* OV588	**3887**	312	1954	**1643**	1955	2242	**288**	2243	2488	**246**	2646	3265	**620**
WP_102076025.1	*Rahnella victoriana* BRK18a	**4027**	312	2030	**1719**	2031	2318	**288**	2319	2564	**246**	2722	3342	**621**
WP_071683781.1	*Serratia fonticola* AeS1	**3994**	298	2000	**1703**	2001	2332	**332**	2333	2517	**185**	2751	3297	**547**
WP_065683710.1	*Serratia* sp. 14-2641	**3848**	297	2000	**1704**	2001	2332	**332**	2333	2517	**185**	2689	3297	**609**
WP_004956787.1	*Serratia odorifera* DSM 4582	**3986**	297	2000	**1704**	2001	2342	**342**	2343	2527	**185**	2752	3307	**556**
WP_051791190.1	*Serratia* sp. DD3	**3350**	335	1484	**1150**	1485	1768	**284**	1769	1953	**185**	2053	2735	**683**
WP_013814704.1	*Serratia plymuthica* AS9	**3314**	419	1464	**1046**	1465	1748	**284**	1749	1933	**185**	2101	2642	**542**
WP_020439873.1	*Serratia plymuthica* S13	**3337**	313	1464	**1152**	1465	1746	**282**	1747	1931	**185**	2098	2714	**617**
WP_012147097.1	***Serratia proteamaculans* 568**	**3040**	313	1164	**852**	1165	1432	**268**	1433	1617	**185**	1784	2400	**617**
WP_017894168.1	*Serratia* sp. S4	**2993**	313	1164	**852**	1165	1432	**268**	1433	1617	**185**	1784	2400	**617**
WP_084297639.1	*Serratia* sp. ATCC 39006	**3690**	300	1812	**1513**	1813	2155	**343**	2156	2341	**186**	2452	3049	**598**
WP_049254954.1	*Shigella boydii* 1221_SBOY	**3213**	309	1398	**1090**	1399	1655	**257**	1656	1840	**185**	2070	2546	**477**
EIQ74285.1	***Shigella flexneri* 1235-66**	**3961**	258	2002	**1745**	2003	2358	**356**	2359	2543	**185**	2576	3320	**745**
WP_051857422.1	*Trabulsiella guamensis* ATCC 49490	**3707**	311	1438	**1128**	1439	1737	**299**	1738	1921	**184**	2102	2800	**699**
WP_054181051.1	*Trabulsiella odontotermitis* TbO1.1	**3189**	309	1262	**954**	1263	1552	**290**	1553	1737	**185**	1908	2516	**609**
WP_038156365.1	*Trabulsiella guamensis* ATCC 49490	**3299**	311	1438	**1128**	1439	1731	**293**	1732	1915	**184**	2099	2696	**598**
EEP92680.1	*Yersinia kristensenii* ATCC 33638	**3396**	337	1484	**1148**	1485	1765	**281**	1766	1950	**185**	2182	2737	**556**
WP_099461935.1	*Yersinia massiliensis* SCPM-O-B-8024	**3311**	368	1484	**1117**	1485	1765	**281**	1766	1950	**185**	2126	2737	**612**
CNE70603.1	*Yersinia kristensenii* 119/84	**3322**	337	1484	**1148**	1485	1765	**281**	1766	1950	**185**	2182	2812	**631**
WP_072087237.1	*Yersinia frederiksenii* IP23698	**3311**	354	1484	**1131**	1485	1765	**281**	1766	1950	**185**	2126	2737	**612**
CQJ01032.1	*Yersinia frederiksenii* 112/02	**2998**	354	1484	**1131**	1485	1765	**281**	1766	1950	**185**	2126	2737	**612**
WP_005186610.1	*Yersinia intermedia* ATCC 29909	**3435**	397	1484	**1088**	1485	1765	**281**	1766	1950	**185**	2128	2737	**610**
WP_050133220.1	*Yersinia frederiksenii* BR166/97	**2957**	515	1074	**560**	1075	1342	**268**	1343	1528	**186**	1705	2311	**607**
WP_050536725.1	*Yersinia mollaretii* 92/84	**2964**	304	1074	**771**	1075	1342	**268**	1343	1528	**186**	1628	2376	**749**
WP_050141948.1	*Yersinia enterocolitica* YE8850	**2964**	307	1074	**768**	1075	1342	**268**	1343	1528	**186**	1628	2376	**749**
WP_004876812.1	***Yersinia mollaretii* ATCC 43969**	**2963**	418	1074	**657**	1075	1342	**268**	1343	1528	**186**	1628	2376	**749**
WP_050535527.1	*Yersinia mollaretii* 61/02	**2993**	307	1074	**768**	1075	1342	**268**	1343	1528	**186**	1628	2376	**749**

[a] The receptor-binding domain (RBD) was determined as the variable region between the last repeat of the FHA-1 region and the YPLP motif of the YP domain.

[b] The FHA-2 region was measured according to annotations on the NCBI server.

Table S3. Oligonucleotides, Related to Figure STAR Methods.

Identifier	Descriptor	Sequence[a]
CH403	T7-BglII	5′ - AAA AGA TCT TAA TAC GAC TCA CTA TAG GAA TTG TGA GCG GAT AAC
CH2770	minE-Nco-for	5′ - TTT CCA TGG CAT TAC TCG ATT TCT TTC TCT C
CH2771	minE-Bam-rev	5′ - TTT GGA TCC GGC TA TTT CAG CTC TTC TG
CH4239	STEC-Kpn-for	5′' - CAG CGG TAC CCT CTC CG
CH4240	STEC-L899-Xho-rev	5′ - TTT CTC GAG CAG AGT GAT GTT CTG CCC C
CH4241	STEC-R1000-Xho-for	5′ - AAA CTC GAG AGA GAC ATC AGC AAT AGT GGG C
CH4243	STEC-E1399-Xho-rev	5′ - TTT CTC GAG CTC AAC CGG AAT GTA TGA TTT TGT C
CH4244	STEC-G1500-Xho-for	5′ - AAA CTC GAG GGT TCA CAA CTG AAT AAT CAG TCC TTC
CH4245	STEC-A1853-Nco/Xho-rev	5′ - TTT CTC GAG CCA TGG CTG CCA GGT GAA CAT TCT GAC
CH4247	STEC-V2323-AfI/Not-rev	5′ - TTT GCG GCC GCC TTA AGG ACC GCA TTA TTG ATG GCA CTG
CH4282	STEC-AfI/Sph-for	5′ - TTT GCA TGC TTA AGG CCG GTG GTA ACA C
CH4283	STEC-S2917-Nhe/Xho-rev	5′ - TTT CTC GAG GCT AGC GCT GCT GTT TCC GG
CH4371	STEC(3)-Nhe/AfI-for	5′ - AAA GCT AGC CTT AAG GCC GGT GGT AAC AC
CH4372	STEC(3)-T2780-Sac-rev	5′ - TTT GAG CTC CGT ATC ACG CAG TTT CGC AAG
CH4373	STEC(3)-L2881-Sac-for	5′ - AAA GAG CTC CTG TAT CCG GGC GTG AAA C
CH4374	STEC(3)-Xho-rev	5′ - TTT CTC GAG TGG CCG GAA TCT
CH4381	STEC(3)-Y1657-Xho-rev	5′ - TTT CTC GAG ATA GTT CCC CTC CAC GAT GC
CH4382	STEC(3)-G1708-Xho-for	5′ - AAA CTC GAG GGG CAG GCA CCA CG
CH4383	STEC(3)-P1707-Xho-rev	5′ - TTT CTC GAG GGG GTG CAT CCT GAG C
CH4384	STEC(3)-Y1758-Xho-for	5′ - AAA CTC GAG TAT ATG CTG AAC CAG ATC GGG G
CH4412	STEC(3)-N2934A-for	5′ - AGA GCG GGA AGA ATG CTG TTG ATA ATG CCT ACC TGA GCG TG
CH4413	STEC(3)-A2918-Nhe-for	5′ - AGC GCT AGC GCA GCA GTG GGC GCA CAG AGC GGG AAG AAT GC
ZR256	tsx-Not-for	5′ - TTT GCG GCC GCG AAT TCG GGA TTT TCA AAC AGT GGC ATA C
ZR257	tsx-Xho-rev	5′ - TTT CTC GAG TCT AGA AAA TCC CGG CAT TTT CAT AAT CAG
ZR260	J2M139 CDI-Not-for	5′ - TTT GCG GCC GCA ATG TCT GGT TGT GGC AGG
ZR261	J2M139 CDI-Xho-rev	5′ - TTT CTC GAG TGG CCG GAA TCT TTA CTC AG
ZR267	cdiA(S504C)(STEC)-Kpn-rev	5′ - GAG GGT ACC GCA GTT GTT TAT CTG CGT GGC AC
ZR268	STEC-A498-Kpn-for	5′ - GCC ACG CAG ATA AAC AAC AGC
ZR269	STEC-cdiA(E512)-rev	5′ - GTT CCT TTG CCC CGG AG
ZR271	cdiA(S1550C)(STEC)-Sac-rev	5′ - CCG GAG CTC CAC AGA GAG TAA AGG TAA CGC GAT CAT C
ZR274	STEC-D1692-Sal-rev	5′ - GCT GTC GAC CTT TCC GAG ACC ATC CAG
ZR275	STEC-S1693C-Sal-for	5′ - AAG GTC GAC TGT AGT CTG TTT GCC GGA CTG TAT GAC
ZR277	STEC-G1726C-Xba-rev	5′ - CCG GTC TAG AAA ATA CGA TGA ACA CGG GAA TTG TTT TTC ATC AGT ATA CG
ZR281	STEC-L1731-Xba-for	5′ - TTT TCT AGA CCG GCT CGG GCT GAA AC
ZR280	STEC-cdiA(A1807C)-Sex-for	5′ - CAG ACC AGG TGG CGT GTC TTG ACC AAA GTA TTC TGT GGT ATA AAG
ZR285	STEC-cdiA(D1959)-rev	5′ - GTC CAG ACT ATC CAT GGC AC
ZR286	cdiA(D1959C)(STEC)-Nco-for	5′ - GTG CCA TGG ATA GTC TGT GCA TCC GGG CAG ATA AAA ACA TTT CG
ZR287	STEC-R1961-Not-rev	5′ - TTT GCG GCC GCG GAT GTC CAG ACT ATC CAT GG
ZR288	STEC-T2502C-Eco-rev	5′ - ACC AGA ATT CAG GGT GAC ACA TTT ACC GCT GTC TGT TGT GG
ZR290	STEC-N2506-Eco-for	5′ - ACC CTG AAT TCT GGT CGG GAT ACG GTA CTG
ZR293	STEC-G2648-Spe-for	5′ - TTT ACT AGT GGC AGC TTC GGG GAT AAA TTT C
ZR294	cdiA(S3217C)(STEC)-Mfe-for	5′ - CGG CAA TTG GTT ATT GCA ATG CAA TAA AAA ATG GAT TAC AGG C
ZR295	malE-Not-for	5′ - TTT GCG GCC GCC AAC AAG GAC CAT AGA TTA TG
ZR296	malE-dSS-Not-for	5′ - TTT GCG GCC GCA AGG ACC ATA GAT TAT GAA AAT CGA AGA AGG TAA ACT GG
ZR297	malE-CysHis6-Xho-rev	5′ - TTT CTC GAG CTA GTG ATG GTG ATG GTG ATG AGA TCC ACC ACA CTT GGT GAT ACG AGT CTG
ZR347	cdiA(A1954)(STEC)-Not-for	5′ - TTT GCG GCC GCC ATG GAT AGT CTG GAC ATC C
ZR348	cdiA(S2731C)(STEC)-Bam-rev	5′ - TTT GGA TCC TAT GTC GCA GAT AAG CCC CAC TGT CTG C
ZR349	cdiA(G2734)(STEC)-Bam-for	5′ - TTT GGA TCC CAG GTG GCG GAT ATC GCG

[a] Underlined residues indicate restriction endonuclease recognition sites.